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RAW SEQUENCE LISTING   DATE: 03/05/2002
PATENT APPLICATION: US/09/899,569A   TIME: 10:07:23

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Output Set: N:\CRF3\03052002\I899569A.raw

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 5       Marwa Scherl-Mostageer
 6       Wolfgang Sommergruber
 7       Roger Abseher
 9 <120> TITLE OF INVENTION: Tumorassoziiertes Antigen (B345)
11 <130> FILE REFERENCE: 0652.2280001
13 <140> CURRENT APPLICATION NUMBER: US/09/899,569A
14 <141> CURRENT FILING DATE: 2001-07-06
16 <150> PRIOR APPLICATION NUMBER: DE 100 33 080.0
17 <151> PRIOR FILING DATE: 2000-07-07
19 <150> PRIOR APPLICATION NUMBER: DE 101 19 294.0
20 <151> PRIOR FILING DATE: 2001-04-19
22 <150> PRIOR APPLICATION NUMBER: US 60/243,158
23 <151> PRIOR FILING DATE: 2000-10-25
25 <150> PRIOR APPLICATION NUMBER: US 60/297,747
26 <151> PRIOR FILING DATE: 2001-06-14
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81 acgagaaagc aacattacag ttctcataaa gctggggacc ccgactctgc tggcaaaacc 180
83 ctgttacatc gtcatttcta aaagacatat aacc atg ttg tcc atc aag tct gga 235
84                                       Met Leu Ser Ile Lys Ser Gly   ·
85                                         1               5
87 gaa aga ata gtc ttt acc ttt agc tgc cag agt cct gag aat cac ttt 283
88 Glu Arg Ile Val Phe Thr Phe Ser Cys Gln Ser Pro Glu Asn His Phe
89         10                  15                  20
91 gtc ata gag atc cag aaa aat att gac tgt atg tca ggc cca tgt cct 331
92 Val Ile Glu Ile Gln Lys Asn Ile Asp Cys Met Ser Gly Pro Cys Pro
93     25                  30                  35
95 ttt ggg gag gtt cag ctt cag ccc tcg aca tcg ttg ttg cct acc ctc 379
96 Phe Gly Glu Val Gln Leu Gln Pro Ser Thr Ser Leu Leu Pro Thr Leu
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100 aac aga act ttc atc tgg gat gtc aaa gct cat aag agc atc ggt tta 427
101 Asn Arg Thr Phe Ile Trp Asp Val Lys Ala His Lys Ser Ile Gly Leu
102                 60                  65                  70
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105 Glu Leu Gln Phe Ser Ile Pro Arg Leu Arg Gln Ile Gly Pro Gly Glu
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109 Ser Cys Pro Asp Gly Val Thr His Ser Ile Ser Gly Arg Ile Asp Ala
110         90                  95                  100
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113 Thr Val Val Arg Ile Gly Thr Phe Cys Ser Asn Gly Thr Val Ser Arg
114     105                 110                 115
116 atc aag atg caa gaa gga gtg aaa atg gcc tta cac ctc cca tgg ttc 619
117 Ile Lys Met Gln Glu Gly Val Lys Met Ala Leu His Leu Pro Trp Phe
118 120                 125                 130                 135
120 cac ccc aga aat gtc tcc ggc ttc agc att gca aac cgc tca tct ata 667
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122                 140                 145                 150
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126                 155                 160                 165
128 acc ctg atg tct gcc aac tac cca gaa ggc ttc cct gag gat gag ctc 763
129 Thr Leu Met Ser Ala Asn Tyr Pro Glu Gly Phe Pro Glu Asp Glu Leu
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141 Glu Tyr Tyr Ile Pro Gly Ser Thr Thr Asn Pro Glu Val Phe Lys Leu
142                    220                    225                    230
144 gag gac aag cag cct ggg aac atg gcg ggg aac ttc aac ctc tct ctg    955
145 Glu Asp Lys Gln Pro Gly Asn Met Ala Gly Asn Phe Asn Leu Ser Leu
146                    235                    240                    245
148 caa ggc tgt gac caa gat gcc caa agt cca ggg atc ctc cgg ctg cag   1003
149 Gln Gly Cys Asp Gln Asp Ala Gln Ser Pro Gly Ile Leu Arg Leu Gln
150            250                    255                    260
152 ttc caa gtt ttg gtc caa cat cca caa aat gaa agc aat aaa atc tac   1051
153 Phe Gln Val Leu Val Gln His Pro Gln Asn Glu Ser Asn Lys Ile Tyr
154        265                    270                    275
158 gtg gtt gac ttg agt aat gag cga gcc atg tca ctc acc atc gag cca   1099
159 Val Val Asp Leu Ser Asn Glu Arg Ala Met Ser Leu Thr Ile Glu Pro
160 280                    285                    290                    295
162 cgg ccc gtc aaa cag agc cgc aag ttt gtc cct ggc tgt ttc gtg tgt   1147
163 Arg Pro Val Lys Gln Ser Arg Lys Phe Val Pro Gly Cys Phe Val Cys
164                    300                    305                    310
167 cta gaa tct cgg acc tgc agt agc aac ctc acc ctg aca tct ggc tcc   1195
168 Leu Glu Ser Arg Thr Cys Ser Ser Asn Leu Thr Leu Thr Ser Gly Ser
169            315                    320                    325
171 aaa cac aaa atc tcc ttc ctt tgt gat gat ctg aca cgt ctg tgg atg   1243
172 Lys His Lys Ile Ser Phe Leu Cys Asp Asp Leu Thr Arg Leu Trp Met
173            330                    335                    340
175 aat gtg gaa aaa acc ata agc tgc aca gac cac cgg tac tgc caa agg   1291
176 Asn Val Glu Lys Thr Ile Ser Cys Thr Asp His Arg Tyr Cys Gln Arg
177        345                    350                    355
179 aaa tcc tac tca ctc cag gtg ccc agt gac atc ctc cac ctg cct gtg   1339
180 Lys Ser Tyr Ser Leu Gln Val Pro Ser Asp Ile Leu His Leu Pro Val
181 360                    365                    370                    375
183 gag ctg cat gac ttc tcc tgg aag ctg ctg gtg ccc aag gac agg ctc   1387
184 Glu Leu His Asp Phe Ser Trp Lys Leu Leu Val Pro Lys Asp Arg Leu
185                    380                    385                    390
187 agc ctg gtg ctg gtg cca gcc cag aag ctg cag cag cat aca cac gag   1435
188 Ser Leu Val Leu Val Pro Ala Gln Lys Leu Gln Gln His Thr His Glu
189            395                    400                    405
191 aag ccc tgc aac acc agc ttc agc tac ctc gtg gcc agt gcc ata ccc   1483
192 Lys Pro Cys Asn Thr Ser Phe Ser Tyr Leu Val Ala Ser Ala Ile Pro
193            410                    415                    420
195 agc cag gac ctg tac ttc ggc tcc ttc tgc ccg gga ggc tct atc aag   1531
196 Ser Gln Asp Leu Tyr Phe Gly Ser Phe Cys Pro Gly Gly Ser Ile Lys
197        425                    430                    435
199 cag atc cag gtg aag cag aac atc tcg gtg acc ctt cgc acc ttt gcc   1579
200 Gln Ile Gln Val Lys Gln Asn Ile Ser Val Thr Leu Arg Thr Phe Ala
201 440                    445                    450                    455
203 ccc agc ttc caa caa gag gcc tcc agg cag ggt ctg acg gtg tcc ttt   1627
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211 aaa agc aag gtc tac ctg agg acc ccc aac tgg gac cgg ggc ctg cca     1723
212 Lys Ser Lys Val Tyr Leu Arg Thr Pro Asn Trp Asp Arg Gly Leu Pro
213         490                     495                 500
216 tcc ctc acc tct gtg tcc tgg aac atc agc gtg ccc aga gac cag gtg     1771
217 Ser Leu Thr Ser Val Ser Trp Asn Ile Ser Val Pro Arg Asp Gln Val
218         505                     510                 515
220 gcc tgc ctg act ttc ttt aag gag cgg agc ggc gtg gtc tgc cag aca     1819
221 Ala Cys Leu Thr Phe Phe Lys Glu Arg Ser Gly Val Val Cys Gln Thr
222 520                 525                 530                 535
224 ggg cgc gca ttc atg atc atc cag gag cag cgg acc cgg gct gag gag     1867
225 Gly Arg Ala Phe Met Ile Ile Gln Glu Gln Arg Thr Arg Ala Glu Glu
226                 540                 545                 550
228 atc ttc agc ctg gac gag gat gtg ctc ccc aag cca agc ttc cac cat     1915
229 Ile Phe Ser Leu Asp Glu Asp Val Leu Pro Lys Pro Ser Phe His His
230                 555                 560                 565
232 cac agc ttc tgg gtc aac atc tct aac tgc agc ccc acg agc ggc aag     1963
233 His Ser Phe Trp Val Asn Ile Ser Asn Cys Ser Pro Thr Ser Gly Lys
234                 570                 575                 580
236 cag cta gac ctg ctc ttc tcg gtg aca ctt acc cca agg act gtg gac     2011
237 Gln Leu Asp Leu Leu Phe Ser Val Thr Leu Thr Pro Arg Thr Val Asp
238         585                 590                 595
240 ttg act gtc atc ctc atc gca gcg gtg gga ggt gga gtc tta ctg ctg     2059
241 Leu Thr Val Ile Leu Ile Ala Ala Val Gly Gly Gly Val Leu Leu Leu
242 600                 605                 610                 615
244 tct gcc ctc ggg ctc atc att tgc tgt gtg aaa aag aag aaa aag aag     2107
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246                 620                 625                 630
248 aca aac aag ggc ccc gct gtg ggt atc tac aat ggc aac atc aat act     2155
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258         665                 670                 675
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262 680                 685                 690                 695
264 acc ggc cgt tcc agg gca cca tgg ggg tct gtc ctc cct ccc cac cca     2347
265 Thr Gly Arg Ser Arg Ala Pro Trp Gly Ser Val Leu Pro Pro His Pro
266                 700                 705                 710
268 cca tat gct cca ggg ccc caa ctg caa agt tgg cca ctg agg agc cac     2395
269 Pro Tyr Ala Pro Gly Pro Gln Leu Gln Ser Trp Pro Leu Arg Ser His
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279 Ile Pro Thr Met Gly Met
280       745
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PATENT APPLICATION:　 US/09/899,569A　　　TIME: 10:07:24

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